

Signal Transduction, Pharmacological Properties, and Expression Patterns of Two Rat Metabotropic Glutamate Receptors, mGluR3 and mGluR4

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The metabotropic glutamate receptors are coupled to intracellular signal transduction via G-proteins and consist of a family of at least five different subtypes, termed mGluR1–mGluR5. We studied the signal transduction mechanism and pharmacological characteristics of the rat mGluR3 and mGluR4 subtypes in Chinese hamster ovary cells permanently expressing the cloned receptors. Both mGluR3 and mGluR4 inhibit the forskolin-stimulated accumulation of intracellular cAMP formation in response to agonist interaction. Consistent with the high degree of sequence similarity to mGluR2, mGluR3 closely resembles mGluR2 in its agonist selectivity; the potency rank order of agonists is L-glutamate > *trans*-1-aminocyclopentane-1,3-dicarboxylate > ibotenate > quisqualate. mGluR4 is totally different in its agonist specificity from any other member of the metabotropic receptors. This receptor potently reacts with L-2-amino-4-phosphonobutyrate (L-AP4) in a stereo-selective manner and moderately responds to L-serine-O-phosphate. mGluR4 thus corresponds well to the putative L-AP4 receptor characterized from brain preparations. Blot and *in situ* hybridization analyses indicated that both mRNAs are widely distributed in the rat brain. mGluR3 mRNA is highly expressed in neuronal cells of the cerebral cortex and the caudate-putamen, and in granule cells of the hippocampal dentate gyrus. The expression pattern of mGluR4 mRNA is more restricted, and this expression is prominent in the cerebellum, olfactory bulb, and thalamus. Furthermore, the mGluR3 mRNA, unlike the other mRNAs for the metabotropic receptors, is highly expressed in glial cells throughout the brain regions. The metabotropic glutamate receptor subtypes can thus be classified into three subgroups according to the similarity in their amino acid sequences, signal transduction, and agonist selectivity: mGluR1/mGluR5, mGluR2/mGluR3, and mGluR4. The mRNAs for the individual receptor subtypes, however, show overlapping but distinct patterns of expression in the rat CNS.

[Key words: metabotropic glutamate receptor, cDNA clone, DNA transfection, signal transduction, agonist selectivity, *in situ* hybridization]

L-Glutamate acts as a major excitatory neurotransmitter and plays an important role in neuronal plasticity and neurotoxicity in the CNS (Monaghan et al., 1989; Watkins et al., 1990). Glutamate neurotransmission is thus thought to be involved in many integrative brain functions including memory acquisition and learning, and also in some neurodegenerative disorders such as stroke and epilepsy (Collingridge and Singer, 1990; Meldrum and Garthwaite, 1990). The disparate functions of glutamate neurotransmission are reflected in the presence of multiple receptors that can be categorized into two distinct groups termed ionotropic and metabotropic receptors (mGluRs) (Monaghan et al., 1989). The ionotropic receptors, consisting of the NMDA receptors and AMPA/kainate receptors, contain cation-specific ion channels and share fundamental features with other ligand-gated ion channels (Hollmann et al., 1989; Monaghan et al., 1989; Moriyoshi et al., 1991). The metabotropic receptors modulate the production of intracellular second messengers and belong to the family of G-protein-coupled receptors (Schoepp et al., 1990; Récasens et al., 1991). An additional and novel receptor termed L-AP4 (L-2-amino-4-phosphonobutyrate) receptor has been identified through the actions of this compound on certain glutamate-using synapses (Monaghan et al., 1989). The L-AP4 receptor has been shown to suppress synaptic glutamate neurotransmission and appears to represent an autoreceptor (Cotman et al., 1986; Forsythe and Clements, 1990; Bas-lys and Malenka, 1991). However, the molecular nature of the L-AP4 receptor remained to be elucidated.

Recently, we isolated cDNA clones for five different subtypes of the rat mGluR family (mGluR1–mGluR5) by molecular screening of a rat brain cDNA library (Masu et al., 1991; Abe et al., 1992; Tanabe et al., 1992). The mGluR family possesses a large extracellular domain preceding the seven putative transmembrane segments and shares a high sequence similarity among the five different subtypes (Houamed et al., 1991; Masu et al., 1991; Abe et al., 1992; Tanabe et al., 1992). These subtypes can be divided further into three subgroups according to their sequence similarities: mGluR1/mGluR5, mGluR2/mGluR3, and mGluR4 (Abe et al., 1992; Tanabe et al., 1992). Consistent with the sequence similarity, both mGluR1 and mGluR5 are coupled to the stimulation of phosphatidylinositol (PI) hydrolysis/Ca²⁺ signal transduction and show strong resemblance in their agonist

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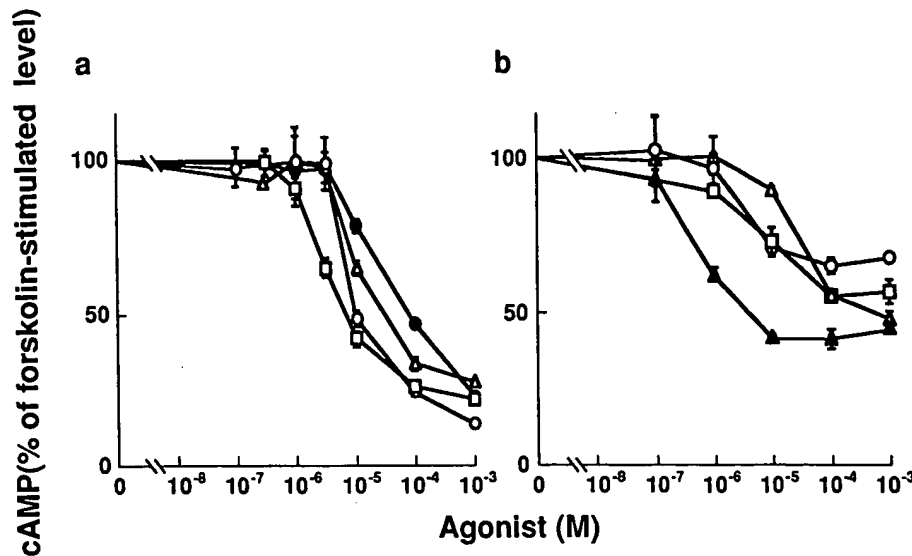


Figure 1. Dose-response curves of agonists in inhibition of forskolin-stimulated cAMP accumulation in cells stably expressing mGluR3 (*a*) and mGluR4 (*b*). Agonists added to forskolin-treated cells are as follows: in *a*, \square , L-glutamate; \circ , tACPD; Δ , ibotenate; \bullet , quisqualate; in *b*, \blacktriangle , L-AP4; \square , L-glutamate; \circ , L-serine-O-phosphate; Δ , D-AP4. Intracellular cAMP levels in cells treated and untreated with $10 \mu\text{M}$ forskolin were 233 ± 8.9 and 7.2 ± 0.4 pmol/well in *a*, respectively, and 239 ± 20 and 8.4 ± 1.0 pmol/well in *b*, respectively. The cAMP levels in forskolin-treated cells are taken as 100%. Each point represents the mean \pm SEM of at least two separate experiments done in triplicate.

selectivity (Abe et al., 1992; Aramori and Nakanishi, 1992). mGluR2, on the other hand, is linked to the inhibitory cAMP cascade and exhibits an agonist selectivity different from that of mGluR1/mGluR5 (Tanabe et al., 1992). However, neither the precise signal transduction nor the agonist selectivity of mGluR3 or mGluR4 has yet been characterized. In this study, we stably expressed mGluR3 and mGluR4 by DNA transfection into Chinese hamster ovary (CHO) cells and investigated the intracellular signal transduction mechanisms and pharmacological characteristics of these receptor subtypes. We also analyzed expression patterns of the mRNAs for mGluR3 and mGluR4 by RNA blot and *in situ* hybridization. The results demonstrated that both subtypes are linked to the inhibitory cAMP cascade. However, mGluR4 is totally different from other members of mGluRs in agonist selectivity and reacts with L-AP4 with a high potency that is one order more effective than L-glutamate. Furthermore, the mGluR3 mRNA, unlike other mRNAs of the mGluR subtypes, has been found to be highly expressed in both neuronal and glial cells.

Materials and Methods

Materials. L-AP4, D-AP4, L-serine-O-phosphate, quisqualate, and tACPD (*trans*-1-aminocyclopentane-1,3-dicarboxylate) were purchased from Tocris Neuramin. D-AP4 was more than 98% enantiomerically pure but may have contained a trace amount of L-AP4 (less than 2%). All other compounds were reagent grade and were obtained as described previously (Tanabe et al., 1992).

Receptor expression in CHO cells and cAMP measurements. The cDNA clones for mGluR3 (pmGR3) and for mGluR4 (pmGR4) were described previously (Tanabe et al., 1992). The AT-rich sequence in the 3' non-coding region of pmGR3 was removed by digestion with PstI, whose site was 65 base pairs (bp) downstream of the stop codon. The resulting 2.9 kilobase pair (kbp) ClaI-PstI fragment of pmGR3 and the 3.9 kbp EcoRI fragment of pmGR4 were inserted individually into a eukaryotic expression vector (pdkCR-dhfr) containing the mouse dihydrofolate reductase gene as a selective marker (Tanabe et al., 1992). These plasmids were transfected into CHO cells deficient in dihydrofolate reductase activity (CHO-dhfr⁻) (Urlaub and Chasin, 1980) by the calcium phosphate method (Graham and van der Eb, 1973). Cell populations expressing mGluR3 or mGluR4 together with dihydrofolate reductase were selected in Dulbecco's modified Eagle's medium lacking L-glutamine, ribonucleosides, and deoxyribonucleosides, and containing a reduced concentration (2 mM) of L-glutamine (Tanabe et al., 1992). Clonal cell lines expressing high levels of mGluR3 or mGluR4 were identified

by measuring the glutamate-induced inhibition of the forskolin-stimulated cAMP accumulation. For the measurement of cAMP levels, a clonal cell line expressing mGluR3 or mGluR4 was seeded individually in 12-well plates at a density of 1.5×10^5 cells/well and grown for 3 d. After 20 min preincubation in phosphate-buffered saline (PBS) containing 1 mM 3-isobutyl-1-methylxanthine (IBMX) at 37°C, the cells were incubated with fresh PBS containing $10 \mu\text{M}$ forskolin, 1 mM IBMX, and test agents for 10 min. The medium was aspirated, and the reaction was stopped by the addition of 5% trichloroacetic acid. cAMP levels were measured by the RIA kit (Amersham). For pertussis toxin (PTX) treatment, cells were preincubated with various concentrations of PTX for 11 hr at 37°C. Each experiment was carried out at least twice in triplicate.

RNA blot and *in situ* hybridization analyses. RNA blot analysis was carried out by using $10 \mu\text{g}$ of total RNAs isolated from various regions of the brain as described previously (Masu et al., 1991). The cDNA probe used was the 877 bp PstI fragment of pmGR3 and the 1263 bp SmaI fragment of pmGR4. *In situ* hybridization was performed as previously described (Masu et al., 1991). Briefly, ³⁵S-labeled antisense riboprobe corresponding to the 1464 bp HincII-XbaI or 648 bp EcoT14I fragment of pmGR3 or the 1230 bp XhoI-PstI or 568 bp PstI fragment of pmGR4 was transcribed and hybridized as described previously (Tanabe et al., 1992). Sections were exposed to β max-film (Amersham) for 2 weeks or dipped in NTB2 emulsion (Kodak) diluted 1:1 with distilled water, developed after a 4 week exposure, and counterstained with cresyl violet. Control hybridization experiments were carried out in adjacent sections by using the same riboprobe in the presence of an excess of unlabeled probe.

Results

Signal transduction and agonist profiles of mGluR3 and mGluR4

To investigate the pharmacological profiles and signal transduction of mGluR3 and mGluR4, we stably expressed individual cDNA clones encoding these receptor subtypes in mammalian CHO cells. A mouse dihydrofolate reductase gene was used as a selective marker gene that allowed receptor-expressing cells to grow in a medium lacking ribonucleosides and deoxyribonucleosides. We also removed L-glutamine and reduced the concentration of L-glutamine in the culture medium to avoid the possible constitutive activation of these receptors by L-glutamine. Through this procedure, we succeeded in obtaining several cell lines stably expressing mGluR3 and mGluR4. Because mGluR3 shows a high degree (~70%) of sequence similarity to mGluR2 that is coupled to the inhibitory cAMP cascade (Tan-

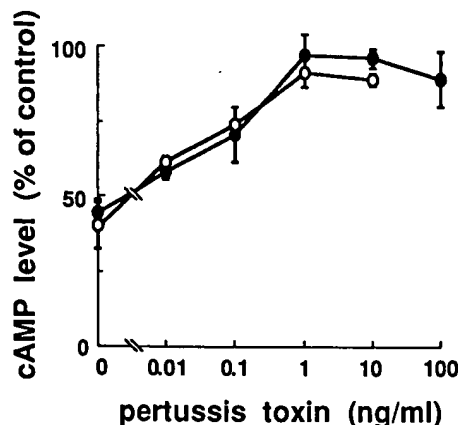


Figure 2. Effects of PTX on agonist-mediated inhibition of forskolin-stimulated cAMP accumulation. mGluR3-expressing cells (○) and mGluR4-expressing cells (●) were pretreated with various concentrations of PTX for 11 hr. Forskolin-stimulated cAMP levels were determined with or without addition of 100 μ M L-glutamate for mGluR3 and with or without addition of 100 μ M L-AP4 for mGluR4. The levels obtained without addition of L-glutamate or L-AP4 are taken as 100% at each concentration of PTX indicated. Each point represents the mean \pm SEM of at least two separate experiments done in triplicate.

abe et al., 1992), we examined inhibitory effects of L-glutamate on the forskolin-stimulated cAMP accumulation in mGluR3-expressing cells. As expected, mGluR3 showed efficient L-glutamate-mediated inhibition of the forskolin-stimulated cAMP formation (Fig. 1*a*). Similarly, L-glutamate added to mGluR4-expressing cells resulted in significant inhibition of the forskolin-induced cAMP accumulation (Fig. 1*b*). This inhibition, however, did not extend beyond about 50% of the maximally stimulated levels even by the addition of higher concentrations of L-glutamate. Because this partial inhibition was consistently observed in several independent mGluR4-expressing cell lines, this inhibition pattern probably represents a feature characteristic of the signal transduction mediated by mGluR4. No L-glutamate-mediated inhibition of the forskolin-stimulated cAMP formation was observed in untransfected control cells or those transfected with the vector DNA alone (data not shown). Furthermore, no appreciable stimulation of PI hydrolysis, cAMP formation, or arachidonic acid release was induced by the addition of L-glutamate in either mGluR3- or mGluR4-expressing cells (data not shown). Thus, it can be concluded that both mGluR3 and mGluR4 subtypes are coupled to the inhibitory cAMP cascade.

We determined dose-response curves of various agonists for the inhibition of the forskolin-stimulated cAMP accumulation in mGluR3-expressing cells (Fig. 1*a*). L-glutamate, tACPD, ibotenate, and quisqualate were effective in inhibiting the cAMP accumulation in this order of agonists. Thus, the rank order of these compounds was in complete agreement with that determined for mGluR2 (Tanabe et al., 1992). The effective concentrations of half-maximal response (EC_{50}) of the above compounds were calculated to be 3, 8, 10, and 40 μ M, respectively. These values were also very similar to those reported for mGluR2 (Tanabe et al., 1992). NMDA, kainate, AMPA, and L-AP4 had virtually no effects on the signal transduction of mGluR3 in receptor-expressing cells. Thus, mGluR3 closely resembles mGluR2 in the signal transduction and agonist selectivity.

Because mGluR4 was found to be linked to the inhibitory cAMP cascade, we determined an agonist profile of mGluR4

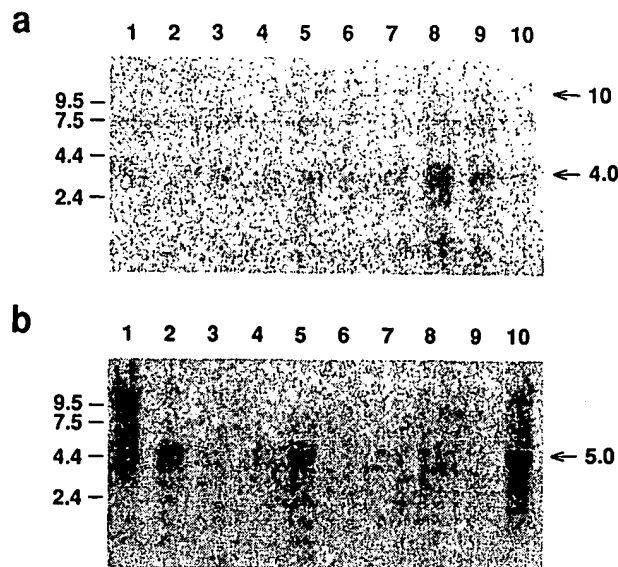


Figure 3. RNA blot hybridization analysis of mGluR3 (*a*) and mGluR4 (*b*). Total RNAs analyzed are as follows: 1, whole brain; 2, olfactory bulb; 3, medulla/pons; 4, midbrain; 5, thalamus; 6, hypothalamus; 7, hippocampus; 8, striatum; 9, cerebral cortex; 10, cerebellum. The size marker (kilobases) used was the RNA ladder (Bethesda Research Labs).

by measuring cAMP levels after application of various compounds to forskolin-treated mGluR4-expressing cells (Fig. 1*b*). The cAMP accumulation was reduced by L-glutamate in a dose-dependent manner with an EC_{50} value of 5 μ M. Notably, L-AP4 more effectively inhibited the forskolin-stimulated cAMP accumulation, and this potency with an EC_{50} value of 0.5 μ M was one order higher than that of L-glutamate. L-Serine-O-phosphate was also an effective agonist with an EC_{50} value of 4 μ M, although the extent of this inhibition was less than those observed for L-AP4 and L-glutamate. D-AP4, an enantiomer of L-AP4, evoked inhibitory responses at higher concentrations. However, the D-AP4 available for our experiments was not entirely pure and may have contained a trace amount of L-AP4. Because L-AP4 and D-AP4 showed about 2 orders of magnitude difference in the effective concentrations, it is feasible that the effect of D-AP4 arises from a small amount of L-AP4 in our D-AP4 preparation rather than reflecting its intrinsic activity. tACPD, quisqualate, and D,L-2-amino-3-phosphonopropionate evoked slight inhibition (10–20%) of the cAMP formation at the concentration of 100 μ M. However, this inhibition was variable among the different experiments, and was also observed in untransfected control cells. Thus, the agonist activity of these compounds could not be evaluated as being statistically significant. NMDA, kainate, ibotenate, and AMPA did not display any inhibitory activity at 100 μ M each. Thus, the results presented here demonstrated that mGluR4 is clearly distinguishable from other members of the mGluR family by its characteristic agonist selectivity.

PTX catalyzes the ADP-ribosylation of some G-proteins, thus uncoupling them from their linked receptors (Gilman, 1984; Ui, 1984). The PTX-sensitive G-proteins include G_i , linked to the inhibitory cAMP cascade, G_o , and a certain type of G_q that may be coupled to PI hydrolysis (Gilman, 1984; Ui, 1984). To address the coupling of mGluR3 and mGluR4 to G-proteins, we investigated the effects of PTX on mGluR3- and mGluR4-me-

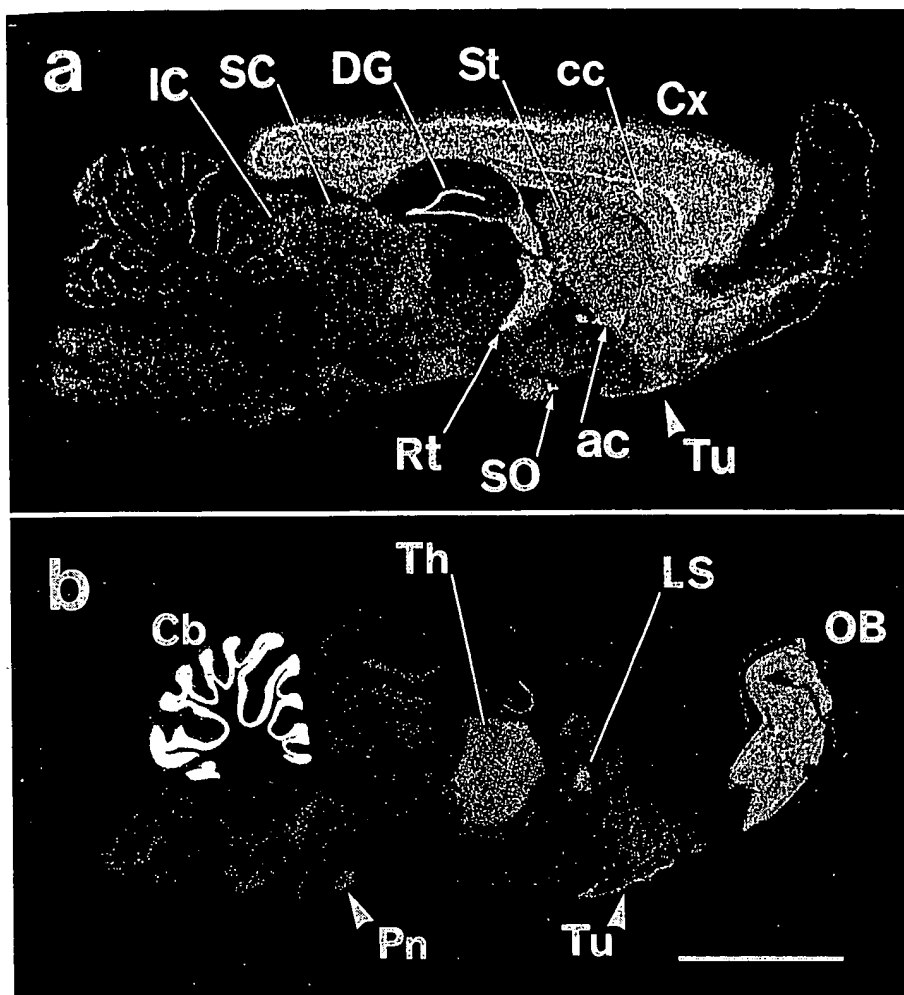


Figure 4. Localization of mGluR3 mRNA (*a*) and mGluR4 mRNA (*b*) in the adult rat brain by *in situ* hybridization. Negative film images of sagittal sections are shown. IC, inferior colliculus; SC, superior colliculus; DG, dentate gyrus; St, striatum; cc, corpus callosum; Cx, cerebral cortex; Rt, thalamic reticular nucleus; SO, supraoptic nucleus; ac, anterior commissure; Tu, olfactory tubercle; Cb, cerebellar cortex; Pn, pontine nucleus; Th, thalamus; LS, lateral septum; OB, main olfactory bulb. Scale bar, 5 mm.

diated inhibition of the cAMP accumulation (Fig. 2). When mGluR3- and mGluR4-expressing cells were pretreated with PTX prior to addition of forskolin and an agonist (L-glutamate for mGluR3 and L-AP4 for mGluR4), the agonist-mediated inhibition of cAMP accumulation was reduced in a dose-dependent manner and was completely abolished at 1 ng of PTX per milliliter of medium in both cases. Thus, the agonist responses in both receptors are mediated by a PTX-sensitive G_i protein.

Expression patterns of mGluR3 and mGluR4 mRNAs

The distribution of mGluR3 and mGluR4 transcripts in the adult rat brain was examined by RNA blot and *in situ* hybridization analyses. Blot hybridization analysis of mGluR3 mRNA revealed a wide distribution of an mRNA of ~4.0 kb throughout the brain regions (Fig. 3*a*). Additionally, a minor band corresponding to an mRNA size of ~10 kb was seen in all regions examined. Blot hybridization analysis of mGluR4 mRNA yielded a band with an estimated mRNA size of ~5.0 kb (Fig. 3*b*). A minor band with an mRNA size of ~10 kb was also found in the cerebellum. The expression pattern of mGluR4 mRNA was rather specific. The high levels of the mRNA expression were observed in the cerebellum, thalamus, and olfactory bulb.

Only a trace amount of the mRNA was seen in the hippocampus.

In situ hybridization analysis indicated that mGluR3 mRNA was expressed not only in neuronal cells but also in glial cells throughout the brain regions (Figs. 4*a*, 5*a-c*). This mRNA was predominantly expressed in neurons of the cerebral cortex, thalamic reticular nucleus (Fig. 5*b*), and supraoptic nucleus, and granule cells in the dentate gyrus, and also in glial cells in white matter such as the corpus callosum (Fig. 5*a*) and anterior commissure. In the cerebellum, moderate expression was seen in Golgi cells (Fig. 5*c*). mGluR4 mRNA was more restrictedly distributed in neuronal cells (Figs. 4*b*, 5*d-f*). Prominent expression of mGluR4 mRNA was observed in neurons of the internal granular layer of the main olfactory bulb (Fig. 5*d*), those of the thalamus (Fig. 5*e*), lateral septum, and pontine nucleus (Fig. 4*b*), and granule cells of the cerebellum (Fig. 5*f*). Only weak expression of mGluR4 mRNA was observed in the dentate gyrus and CA3 region of the hippocampus. No significant hybridization was observed in parallel control experiments using the same mGluR3 or mGluR4 probe in the presence of an excess of unlabeled probe (data not shown). Furthermore, in both mRNA analyses, a similar hybridization pattern was obtained with a different nonoverlapping probe derived from the mGluR3 or mGluR4 cDNA (data not shown).

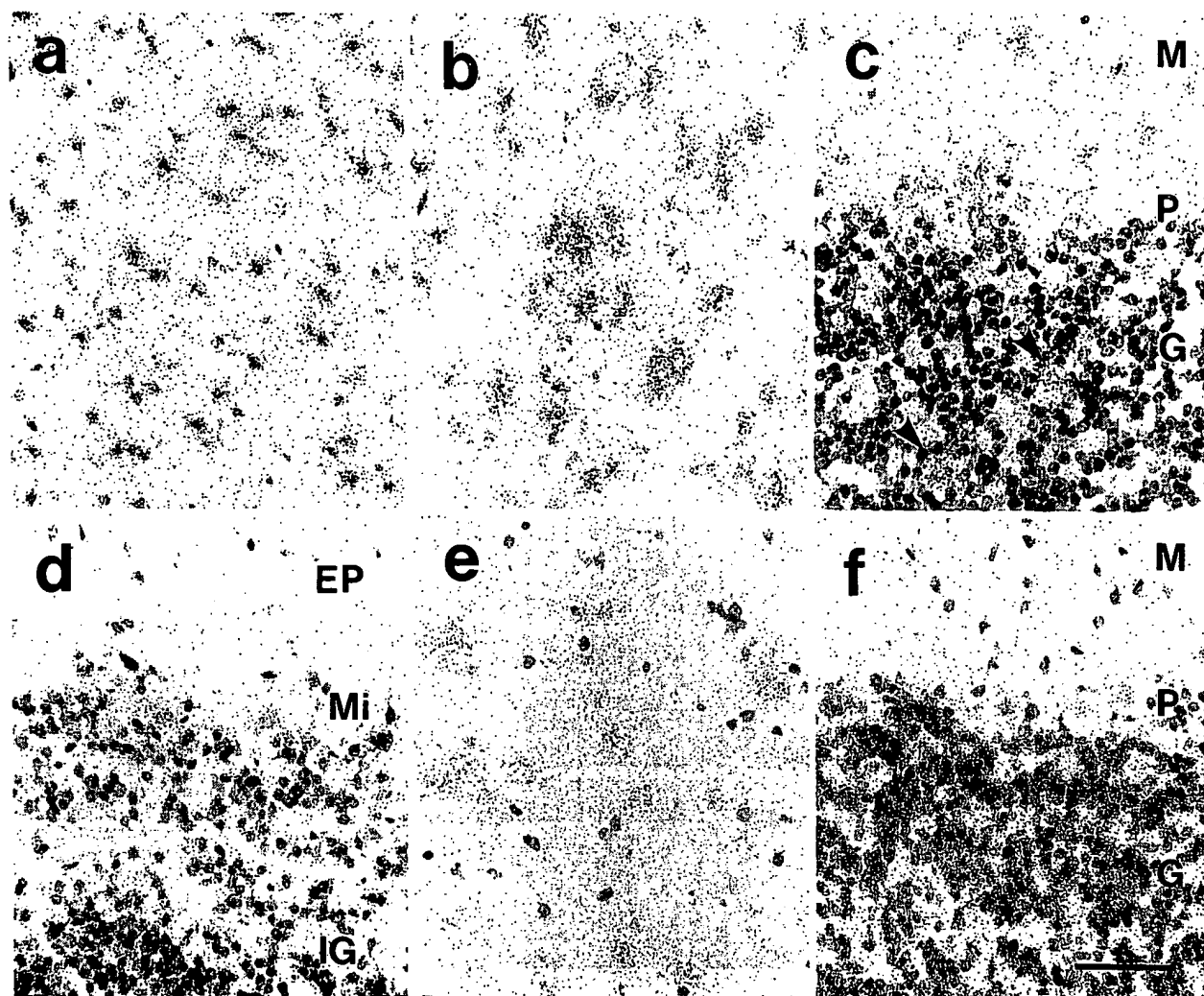


Figure 5. Localization of mGluR3 mRNA (*a–c*) and mGluR4 mRNA (*d–f*) in the adult rat brain by *in situ* hybridization. Bright-field photographs of emulsion-dipped sections through the corpus callosum (*a*), thalamic reticular nucleus (*b*), cerebellar cortex (*c, f*), olfactory bulb (*d*), and ventrolateral thalamic nucleus (*e*) are shown. In *a*, intense signals are seen in small glial cells. Arrowheads in *c* indicate labeled Golgi cells. *M*, molecular layer; *P*, Purkinje cell layer; *G*, granular layer; *EP*, external plexiform layer; *Mi*, mitral cell layer; *IG*, internal granular layer. Scale bar, 50 μ m.

Discussion

This article reports the characterization of the signal transduction and agonist profiles of mGluR3 and mGluR4 and their expression patterns in the adult rat brain. mGluR3 resembles mGluR2 in both signal transduction and agonist selectivity. Similar to mGluR3, mGluR4 is coupled to the inhibitory cAMP cascade, but its agonist selectivity is totally different from not only mGluR2/mGluR3 but also mGluR1/mGluR5. Interestingly, mGluR4 effectively and selectively interacts with L-AP4. It has previously been suggested by electrophysiological studies that there is a novel receptor that potently responds to L-AP4 (Koerner et al., 1981; Hori et al., 1982; Yamamoto et al., 1983; Lanthorn et al., 1984; Hearn et al., 1986). These studies indicated that L-AP4 suppressed neurotransmission in various synapses of the rat CNS, including the lateral entorhinal–dentate gyrus pathway and the lateral olfactory tract–prepyriform cortex pathway. L-AP4-sensitive responses have also been reported to

occur in the mossy fiber–CA3 synapses of guinea pig, but not in the same synapses of rat (Lanthorn et al., 1984). The expression of mGluR4 mRNA is not necessarily high in neuronal cells projecting these neuronal pathways. However, the putative L-AP4 receptor has been shown to exhibit high stereospecificity for the L-isomer of AP4 and to interact moderately with L-serine-O-phosphate (Cotman et al., 1986). Thus, the agonist profile of this receptor agrees well with that of mGluR4. Recent electrophysiological studies indicated that the L-AP4 receptor is involved in suppression of excitatory synaptic transmission by reducing glutamate release at a presynaptic site (Cotman et al., 1986; Forsythe and Clements, 1990; Baskys and Malenka, 1991). Thus, the L-AP4 receptor appears to represent a presynaptic autoreceptor. It is thus feasible that mGluR4 corresponds to the L-AP4 receptor at least at some of neuronal synapses and regulates glutamate neurotransmission by the action at presynaptic sites.

Another interesting system in which L-AP4 has been exten-

Table 1. Properties and expressions of the five subtypes of the mGluR family

Subgroup	Receptor (amino acids)	Signal transduction	Agonist selectivity	mRNA expression sites*
I	mGluR1 (1199 or 906)	IP ₃ /Ca ²⁺ cascade	QA > Glu ≥ Ibo > tACPD	Purkinje cells of cerebellum, mitral and tufted cells of olfactory bulb, granule cells of dentate gyrus, pyramidal cells of CA2–CA4, neurons of thalamus
	mGluR5 (1171)			Neurons of striatum, cerebral cortex, and internal granular layer of olfactory bulb; granule cells of dentate gyrus; pyramidal cells of CA1–CA4
II	mGluR2 (872)	Inhibitory cAMP cascade	Glu ≥ tACPD > Ibo > QA	Golgi cells of cerebellum, granule cells of dentate gyrus, neurons of cerebral cortex and main and accessory olfactory bulb
	mGluR3 (879)			Neurons of cerebral cortex and thalamic reticular nucleus, granule cells of dentate gyrus, glial cells throughout the CNS
III	mGluR4 (912)	Inhibitory cAMP cascade	L-AP4 > Glu > SOP	Granule cells of cerebellum, neurons of thalamus and internal granular layer of olfactory bulb

IP₃, inositol trisphosphate; QA, quisqualate; Glu, glutamate; Ibo, ibotenate; SOP, L-serine-O-phosphate.

* Only characteristic expression sites are described.

sively studied is within the retina. L-glutamate hyperpolarizes ON-bipolar cells and depolarizes OFF-bipolar cells in the retina. L-AP4 mimics the action of L-glutamate at ON-bipolar cells and selectively hyperpolarizes these cells (Nawy and Jahr, 1990, 1991). The hyperpolarization has been shown to result from lowering intracellular cGMP levels through the activation of G-protein-coupled glutamate receptor. The slight difference, however, has been reported for the pharmacological properties of the putative L-AP4 receptors in the hippocampal and retinal preparations (Peterson et al., 1991). It will thus be interesting to investigate whether mGluR4 or its related possible subtype is involved in hyperpolarization of ON-bipolar cells, and whether this receptor subtype is coupled to the cGMP cascade to regulate glutamate neurotransmission in bipolar cells.

Consistent with the sequence conservation between mGluR2 and mGluR3, both receptors closely resemble each other in the signal transduction and agonist selectivity. Recently, tACPD, which is a potent agonist for mGluR2/mGluR3, has been shown to inhibit cAMP formation in slices of the rat hippocampus (Schoepp et al., 1992). Furthermore, this compound showed suppressive effects on excitatory synaptic neurotransmission by the action at a presynaptic site (Baskys and Malenka, 1991). Thus, this receptor subgroup may also serve as a presynaptic autoreceptor. The mGluR3 subtype is, however, distinguished from other members of mGluRs by its prominent expression in glial cells throughout the brain regions. It has been reported that glial cells contain an mGluR that is linked to PI hydrolysis and shows an agonist selectivity different from that of mGluR3 (Pearce et al., 1986; Nicoletti et al., 1990). The discovery of the inhibitory cAMP cascade-linked mGluR in glial cells was un-

expected, and the role of this receptor in glial functions will be interesting for further investigations.

Taking into account the results presented here, together with those reported previously (Abe et al., 1992; Aramori and Nakanishi, 1992; Tanabe et al., 1992), Table 1 summarizes the properties and expressions of five subtypes of the mGluR family. The five receptors can be classified into three subgroups. The first subgroup consists of mGluR1 and mGluR5 and is coupled to PI hydrolysis/Ca²⁺ signal transduction. The most potent agonist for this receptor subgroup is quisqualate. The second subgroup is composed of mGluR2 and mGluR3 and is linked to the inhibitory cAMP cascade. tACPD is a very potent agonist for this subgroup. mGluR4 is classified as the third subgroup and is highly reactive with L-AP4. Although mGluR4 also inhibits cAMP accumulation, its low efficacy in inhibiting the cAMP formation may suggest that mGluR4 has yet another coupling system. The subtypes of the same subgroup show 60–70% identity in the amino acid sequences and about 40% homology with different subgroups of the receptors. The properties characteristic of each subgroup are reflected by the high sequence similarity within the same subgroup of the receptors. Individual receptor subtypes, however, show specialized expression patterns in the CNS. Different subtypes of mGluRs may thus play differential roles in glutamate neurotransmission through the coupling to distinct intracellular signal transduction and different expressions in the CNS.

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